# Science Immunology

# Supplementary Materials for

# Identification of resident memory CD8<sup>+</sup> T cells with functional specificity for SARS-CoV-2 in unexposed oropharyngeal lymphoid tissue

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Table S5



# Figure S1. Flow cytometric gating strategy for the identification of mCD4 $^{+}$ and mCD8 $^{+}$ T

cells.

Numbers indicate percentages in the drawn gates.



Figure S2. Assessment of activation-induced markers and cytokines for the detection of virus-specific tonsillar mCD4<sup>+</sup> and mCD8<sup>+</sup> T cells.

Tonsil cells were stimulated with the indicated peptides or SEB and analyzed for surface expression of activation-induced markers (AIMs) or intracellular expression of AIMs and cytokines (ICS). Negative control wells contained equivalent DMSO. (**A**, **B**) Representative

plots showing the expression profile of surface AIMs (top) and intracellular AIMs/cytokines (bottom) for tonsillar mCD8<sup>+</sup> (A) or mCD4<sup>+</sup> T cells (B). (**C**) Frequencies of mCD8<sup>+</sup> (top) or mCD4<sup>+</sup> T cells (bottom) expressing the indicated markers in the absence of stimulation. (**D**) Net frequencies (background subtracted) of virus-specific mCD8<sup>+</sup> (top) or mCD4<sup>+</sup> T cells (bottom) identified by different marker combinations. (**E**) Stimulation indices for virus-specific mCD8<sup>+</sup> (top) or mCD4<sup>+</sup> T cells (bottom) calculated as fold change relative to the negative control. The dotted line indicates a stimulation index of 2. (C to E) Symbols represent different individuals (n = 2).



#### Figure S3. Evaluation of SARS-CoV-2-specific mCD8<sup>+</sup> T cell responses

(A) Overall frequency of tonsil samples tested with individual peptide pools. n = 81. (B) Overall number of antigens recognized in the same tonsil sample pooled for all donors with positive mCD8<sup>+</sup> T cell responses against at least one antigen derived from SARS-CoV-2. n = 26 (only tonsil samples, for which all 6 SARS-CoV-2 antigens were tested, are included).



Figure S4. Identification of tonsillar mCD4<sup>+</sup> T cells specific for SARS-CoV-2 in unexposed individuals.

Tonsil cells were stimulated with the indicated peptides and analyzed for intracellular coexpression of CD40L and TNF- $\alpha$ . (**A**) Representative plots showing the gating of CD40L<sup>+</sup>TNF- $\alpha$ <sup>+</sup> mCD4<sup>+</sup> T cells from individuals with low (top) or high background levels of activation (bottom). (**B**) Frequencies of CD40L<sup>+</sup>TNF- $\alpha$ <sup>+</sup> mCD4<sup>+</sup> T cells. (**C**) Net frequencies (background subtracted) of CD40L<sup>+</sup>TNF- $\alpha$ <sup>+</sup> mCD4<sup>+</sup> T cells. (**D**) Stimulation indices for

CD40L<sup>+</sup>TNF- $\alpha^+$  mCD4<sup>+</sup> T cells calculated as fold change relative to the negative control. The dotted line indicates a stimulation index of 2. (**E**) Overall frequency of tonsil samples with positive mCD4<sup>+</sup> T cell responses (stimulation index ≥2) to the indicated antigens. (**F**) Overall frequency of tonsil samples with positive mCD4<sup>+</sup> and/or mCD8<sup>+</sup> T cell responses (stimulation index ≥2) to SARS-CoV-2. (**G**) Frequencies of CD40L<sup>+</sup>TNF- $\alpha^+$  mCD4<sup>+</sup> T cells in matched blood and tonsil samples. (B–E) S, spike; NC, nucleocapsid; M+E, membrane+envelope. (B–D) Graphs show median ± IQR. Kruskal-Wallis test with Dunn's posttest. (G) Wilcoxon signed rank test. \**P* < 0.05; \*\**P* < 0.01; \*\*\**P* < 0.001; \*\*\*\**P* < 0.0001.



Figure S5. Evaluation of SARS-CoV-2-specific mCD8<sup>+</sup> T cell responses and HCoV antibody titers in children and adults.

(A) Titers for different SARS-CoV-2-spike-specific antibodies measured by ProcartaPlex assay in unexposed children (n=21), adults (n=12) or SARS-CoV-2 convalescent individuals (n=2). Dotted line represents threshold of seropositivity calculated based on SARS-CoV-2 spike trimer-specific antibody values in unexposed individuals (mean + (3 \* SD)). This threshold was applied to Ig measurements shown in Fig. 2D. (B) Stimulation indices for HCoV-OC43-specific mCD4<sup>+</sup> (left) and mCD8<sup>+</sup> (right) T cell responses in HCoV-OC43 seronegative or seropositive individuals. Dotted line represents stimulation index of 2. (C) Associations between 4-1BB<sup>+</sup>IFN- $\gamma^+$  mCD8<sup>+</sup> T cell stimulation indices for the indicated SARS-CoV-2 antigens and indicated HCoV antibody titers (U/mI) or total mCD8<sup>+</sup> T cell frequencies with corresponding Spearman R and *P* values. (D) Stimulation indices for all SARS-CoV-2-specific mCD8<sup>+</sup> T cell responses in indicated HCoV-seronegative or -seropositive children. Dotted line represents stimulation indices for all SARS-CoV-2-specific mCD8<sup>+</sup> T cell responses in indicated HCoV-seronegative or -seropositive children. Dotted line represents stimulation indices for all SARS-CoV-2-specific mCD8<sup>+</sup> T cell responses in indicated HCoV-seronegative or -seropositive children. Dotted line represents stimulation index of 2. (A) S, spike; NC, nucleocapsid; RBD, receptor binding

domain. (A, B, D) Graphs show median ± IQR. (C) Spearman correlations. (D) Mann-Whitney test. \*P < 0.05; \*\*P < 0.01.



Figure S6. Comparability and stability of the tonsillar mCD8<sup>+</sup> T cell phenotype.

(**A**) Representative plots showing intracellular expression of IFN- $\gamma$  and surface expression of CD69 among mCD8<sup>+</sup> T cells. Surface staining for CD69 was performed either before (top) or after stimulation (bottom). (**B**) Representative plot showing EBV tetramer staining of tonsillar mCD8<sup>+</sup> T cells from one individual. (**C**) Frequencies of CXCR5<sup>+</sup>, CD69<sup>+</sup>, or CD103<sup>+</sup> EBV-specific mCD8<sup>+</sup> T cells identified as tetramer<sup>+</sup> or 4-1BB<sup>+</sup>IFN- $\gamma^+$  (ICS). Symbols represent

different individuals (n=2). (**D**, **E**) Expression profiles for CD69/CD103 (D) or CXCR5 (E) among SARS-CoV-2-reactive mCD8<sup>+</sup> T cells specific for the indicated antigens. (**F**, **G**) Representative plots showing expression of CD69/CD103 (F) or CXCR5 (G) among total (left) or SARS-CoV-2 spike-specific mCD8<sup>+</sup> T cells (right) from one individual. (**H**, **I**) Expression profiles for CD69/CD103 (H) or CXCR5 (I) among virus-specific mCD8<sup>+</sup> T cells from children or adults. (D, E) S, spike; NC, nucleocapsid; M+E, membrane+envelope. (C) Graphs show median. (D, E, H, I) Graphs show median ± IQR. (D, E) Kruskal-Wallis test with Dunn's posttest. (H, I) Mann-Whitney test for comparisons between children and adults, Kruskal-Wallis test with Dunn's posttest for comparisons between antigens. \**P* < 0.05; \*\**P* < 0.01; \*\*\*\**P* < 0.001; ns, not significant.



Figure S7. Functional profiles of virus-specific tonsillar mCD8<sup>+</sup> T cells.

(A) Coexpression of CD107a, TNF- $\alpha$ , and IL-2 among 4-1BB<sup>+</sup>IFN- $\gamma^+$  virus-specific mCD8<sup>+</sup> T cells. (B) Geometric mean (GeoMean) fluorescence values for the same functions in children and adults. (C) Coexpression of CD107a, TNF- $\alpha$ , and IL-2 among 4-1BB<sup>+</sup>IFN- $\gamma^+$  virus-specific mCD8<sup>+</sup> T cells from children or adults. (A-C) Graphs show show median ± IQR. (A) Kruskal-Wallis test with Dunn's post-test. (B) Kruskal-Wallis test with Dunn's post-test for comparisons among virus specificities within each age group; Mann-Whitney test for comparisons between children and adults within virus specificities. (C) Mann-Whitney test. \**P* < 0.05; \*\**P* < 0.01; \*\*\*\**P* < 0.001; \*\*\*\**P* < 0.0001; ns, not significant.

 Table S1. Donor characteristics.

Tonsil cohorts, SARS-CoV-2 unexposed	Children (n = 40)	Adults (n = 41)
Age (median [range], years)	3 [2–5]	41 [28–67]
Sex (male, female)	23, 17	37, 4
Year of surgery	2015–2016	2016–2018

#### **SARS-CoV-2 convalescent** *n* = 2

Age (years)	32, 35
Sex (male, female)	2, 0
Year of collection	2021

	Seropositive children		Seropositive adults	
	Number	Percentage	Number	Percentage
HCoV-229E	4/21	19.1	12/12	100
HCoV-NL63	13/21	62.0	12/12	100
HCoV-HKU1	9/21	42.9	12/12	100
HCoV-OC43	10/21	47.6	12/12	100
Any HCoV	18/21	85.7	12/12	100

# Table S2. Seroprevalence for HCoVs in children and adults

Table S3. Associations between mCD8<sup>+</sup> T cell responses specific for SARS-CoV-2 and mCD8<sup>+</sup> T cell responses specific for EBV, CMV, or HCoV-OC43.

Children		Neg	Pos	Total	P *
EBV	Neg Pos	7 1	0 6	7 7	0.0047
CMV	Neg Pos	6 2	1 3	7 5	0.2222
OC43	Neg Pos	6 0	5 0	11 0	NA
Adults		Neg	Pos	Total	P *
EBV	Neg Pos	2 11	0 10	2 21	0.4862
CMV	Neg Pos	6 6	3 7	9 13	0.4149
OC43					0.1984

\*Fisher's exact test. NA, not applicable.

	Reagent	Catalog number	Source	RRID
	Anti-CCR4–BB700 (clone 1G1)	566475	<b>BD Biosciences</b>	RRID:AB_2744302
	Anti-CCR6–BUV737 (clone 11A9)	612780	<b>BD Biosciences</b>	RRID:AB_2870109
	Anti-CCR7–APC-Cy7 (clone G043H7)	353212	BioLegend	RRID:AB_10916390
	Anti-CD3–BUV805 (clone UCHT1)	612895	<b>BD Biosciences</b>	RRID:AB_2870183
	Anti-CD3–BV605 (clone UCHT1)	300460	BioLegend	RRID:AB_2564380
	Anti-CD4–BUV496 (clone SK3)	612936	<b>BD Biosciences</b>	RRID:AB_2870220
	Anti-CD4–PE-Cy5.5 (clone S3.5)	MHCD0418	Thermo Fisher Scientific	RRID:AB_10376013
	Anti-CD8–BUV395 (clone RPA-T8)	563795	BD Biosciences	RRID:AB_2722501
	Anti-CD14–BV510 (clone M5E2)	301842	BioLegend	RRID:AB_2561946
	Anti-CD19–BV510 (clone HIB19)	302242	BioLegend	RRID:AB_2561668
	Anti-CD45RA–BV570 (clone HI100)	304132	BioLegend	RRID:AB_2563813
	Anti-CD49a–BUV615 (clone SR84)	751437	BD Biosciences	RRID:AB_2875433
	Anti-CD69–BUV563 (clone FN50)	748764	<b>BD Biosciences</b>	RRID:AB_2873167
	Anti-CD103–BV605 (clone Ber-ACT8)	350218	BioLegend	RRID:AB_2564283
Antibodies	Anti-CD107a–BV785 (clone H4A3)	328644	BioLegend	RRID:AB_2565967
	Anti-CD134–PE (clone Ber-ACT35)	350004	BioLegend	RRID:AB_10645478
	Anti-CD137–PE-Cy7 (clone 4B4-1)	309818	BioLegend	RRID:AB_2207741
	Anti-CD154–BV421 (clone 24-31)	310824	BioLegend	RRID:AB_2562721
	Anti-CXCR3–BV750 (clone 1C6)	746895	<b>BD Biosciences</b>	RRID:AB_2871692
	Anti-CXCR5–BB515 (clone RF8B2)	564624	<b>BD Biosciences</b>	RRID:AB_2738871
	Anti-IFN-γ–FITC (clone B27)	554700	<b>BD Biosciences</b>	RRID:AB_395517
	Anti-IFN-γ–PE (clone B27)	506507	BD Biosciences	RRID:AB_315440
	Anti-IL-2–PE-Dazzle594 (clone MQ1-17H12)	500344	BioLegend	RRID:AB_2564091
	Anti-IL-17A–eFluor660 (clone eBio64CAP17)	50-7178-42	Thermo Fisher Scientific	RRID:AB_2574282
	Anti-PD-1–BV711 (clone EH12.2H7)	329928	BioLegend	RRID:AB_2562911
	Anti-PD-L1–BV786 (clone 29E.2A3)	329735	BioLegend	RRID:AB_2629581
	Anti-TNF-α–BV650 (clone Mab11)	502936	<b>BD Biosciences</b>	RRID:AB_2563884
Other reagents	BD GolgiStop	554724	<b>BD Biosciences</b>	-
	Brefeldin A	420601	BioLegend	-
	Brilliant Stain Buffer Plus	566385	BD Biosciences	RRID:AB_2869761
	Dasatinib	73082	STEMCELL	-
	DNase I	4716728001	Sigma-Aldrich	-
	FoxP3 Transcription Factor Staining Buffer Set	00-5523-00	Thermo Fisher Scientific	-
	LIVE/DEAD Fixable Aqua Dead Cell Stain Kit	L34957	Thermo Fisher Scientific	-
	Paraformaldehyde	22023	Biotium	-
	Staphylococcal enterotoxin B (SEB)	S4881	Sigma-Aldrich	-
Software	FlowJo	-	FlowJo, LLC	RRID:SCR_008520
	Prism	-	GraphPad Software Inc.	RRID:SCR_002798
	Bioplex Manager	-	Biorad	RRID:SCR_014330

# Table S4. Flow cytometry reagents and analysis software.